

## **RAW SEQUENCE LISTING**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number:	10/510,876
Source:	PCT
Date Processed by STIC:	07-19-2005

## ENTERED



PCT

RAW SEQUENCE LISTING DATE: 07/19/2005
PATENT APPLICATION: US/10/510,876 TIME: 08:55:04

Input Set : A:\PTO.SR.txt

Output Set: N:\CRF4\07192005\J510876.raw

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3 <110> APPLICANT: Power, Christine
              Plater-Zyberk, Christine
      6 <120> TITLE OF INVENTION: Use of osteoprotegerin for the treatment and/or prevention
of fibrotic
              disease
      9 <130> FILE REFERENCE: SLII-P01-001
     11 <140> CURRENT APPLICATION NUMBER: US 10/510,876
C--> 12 <141> CURRENT FILING DATE: 2004-10-08
     14 <150> PRIOR APPLICATION NUMBER: EP02100364.5
    15 <151> PRIOR FILING DATE: 2002-04-10
     18 <160> NUMBER OF SEQ ID NOS: 12
     20 <170> SOFTWARE: PatentIn version 3.1
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     24 <212> TYPE: DNA
     25 <213> ORGANISM: Homo sapiens
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                                                                              120
     32 cactttacaa gtcatcaagt ctaacttcta gaccagggaa ttaatggggg agacagcgaa
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     34 ccctagagca aagtgccaaa cttctgtcga tagcttgagg ctagtggaaa gacctcgagg
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     36 aggctactcc agaagttcag cgcgtaggaa gctccgatac caatagccct ttgatgatgg
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     38 tggggttggt gaagggaaca gtgctccgca aggttatccc tgccccaggc agtccaattt
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    46 aagaggggcc ctgtaatttg aggtttcaga acccgaagtg aaggggtcag gcagccgggt
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    50 tgcgtccgga tcttggctgg atcggactct cagggtggag gagacacaag cacagcagct
    52 gcccagcgtg tgcccagccc tcccaccgct ggtcccggct gccaggaggc tggccgctgg
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    56 ccggtggctt ttttttcccc tgctctccca ggggacagac accaccgccc caccctcac
    58 gccccacctc cctgggggat cctttccgcc ccagccctga aagcgttaat cctggagctt
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                                                                             1020
    60 totgcacaco cocogacogo tocogoccaa gottoctaaa aaagaaaggt gcaaagtttg
    62 gtccaggata gaaaaatgac tgatcaaagg caggcgatac ttcctgttgc cgggacgcta
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    64 tatataacgt gatgagegea egggetgegg agaegeaceg gagegetege eeageegeeg
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    68 tggtaagtcc ctgggccagc cgacgggtgc ccggcgcctg gggaggctgc tgccacctgg
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    70 teteceaace teecagegga eeggegggga gaaggeteea etegeteeet eecaggagag
    72 gcttggggtt aggctggagc aggaaaccgc tttcaagtta tgccatgctt cccctagggt
                                                                             1380
    74 gtccttttac gctgcaaagt tcctgctgac tttatggaag acagcaagag agagacagac
                                                                             1440
    76 agcgagagag agggagagag agagagagag aaacttgttt gaaagtttta gtcattaacc
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    78 ttctgtcttc atctcagaat attaacgccc tcatgtagtc catactatct ttgcttaatg
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1620

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98 Lys Trp Thr Thr Gln Glu Thr Phe Pro Pro Lys Tyr Leu His Tyr Asp 99 20 25 30 102 Glu Glu Thr Ser His Gln Leu Leu Cys Asp Lys Cys Pro Pro Gly Thr																	
102	Glu	Glu	Thr	Ser	His	Gln	Leu	Leu	Cys	Asp	Lys	Cys	Pro	Pro	Gly	Thr	
103			35					40					45		_		
	Tyr		Lys	Gln	His	Cys		Ala	Lys	Trp	Lys	Thr	Val	Cys	Ala	Pro	
107		50					55					60					
		Pro	Asp	His	Tyr		Thr	Asp	Ser	Trp		Thr	Ser	Asp	Glu		
111		-	_	•	_	70	_	_	<b>~</b> 1	_	75	_		_	~ .	80	
	Leu	Tyr	Cys	Ser		val	Cys	Lys	GIu		Gin	Tyr	vaı	rys		GLu	
115	Cvc	Λan	7 ~~	Th≻	85 Uic	7 00	7. ~~~	W-1	C	90	C	T	C1.,	C1	95 7~~	т	
119	Cys	ASII	ALG	100	птэ	ASII	ALG	Val	105	GIU	Cys	гу	GIU	110	ALG	ıyı	
	I.e.ii	Glu	Tle		Phe	Cvs	T.e.11	Lys		Δrα	Ser	Cvs	Pro		Glv	Phe	
123	Lou	014	115	OI u	1110	Cyb	шец	120		9	001	Cyb	125	110	Ory	1110	
	Glv	Val		Gln	Ala	Glv	Thr	Pro	Glu	Ara	Asn	Thr		Cvs	Lvs	Ara	
127	_	130				- 2	135			5		140	-	- 2 -			
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131	145		=	_		150					155		_			160	
134	Arg	Lys	His	Thr	Asn	Cys	Ser	Val	Phe	Gly	Leu	Leu	Leu	Thr	Gln	Lys	
135					165					170					175		
	Gly	Asn	Ala		His	Asp	Asn	Ile		Ser	Gly	Asn	Ser	Glu	Ser	Thr	
139		_	_	180					185					190			
	Gln	Lys	_	Gly	Ile	Asp	Val	Thr	Leu	Cys	Glu	Glu		Phe	Phe	Arg	
143	<b>D</b> 1	~ 7	195	-	m1	-	<b>5</b> 1	200	_	_			205		_		
	Pne		vaı	Pro	Thr	ьys		Thr	Pro	Asn	Trp		Ser	vaı	ьеи	Val	
147	7 cn	210	Ton	Dro	C1	Th∽	215	Val	7 on	71.7	Clu	220	Wal	C1,,	7.~~	т1 о	
	225	AŞII	ьец	FIQ	сту	230	ту	val	ASII	Ата	235	ser	vaı	GIU	Arg	240	
		Δra	Gln	His	Ser		Gln	Glu	Gln	Thr		Gln	Len	T.011	Luc		
155	_	AL G	GIII	1113	245	261	GIII	GIU	GIII	250	rne	GIII	пец	пеп	255	теи	
			His	Gln		Lvs	Asp	Gln	Asp		Val	Lvs	Lvs	Tle		Gln	
159		-1-		260		_,_			265			-10	-10	270		<b></b>	
	Asp	Ile	Asp	Leu	Cys	Glu	Asn	Ser		Gln	Arq	His	Ile		His	Ala	
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166	Asn	Leu	Thr	Phe	Glu	Gln	Leu	Arg	Ser	Leu	Met	Glu	Ser	Leu	Pro	Gly	
167		290					295	=				300				_	
		Lys	Val	Gly	Ala	Glu	Asp	Ile	Glu	Lys	Thr	Ile	Lys	Ala	Cys	Lys	
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178 Gly Asp Gln Asp Thr Leu Lys Gly Leu Met His Ala Leu Lys His Ser
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                                    345
182 Lys Thr Tyr His Phe Pro Lys Thr Val Thr Gln Ser Leu Lys Lys Thr
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                                360
186 Ile Arg Phe Leu His Ser Phe Thr Met Tyr Lys Leu Tyr Gln Lys Leu
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190 Phe Leu Glu Met Ile Gly Asn Gln Val Gln Ser Val Lys Ile Ser Cys
191 385
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208 cgtgtttctg gacateteca ttaagtggae cacceaggaa acgttteete caaagtaeet
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210 tcattatgac gaagaaacct ctcatcagct gttgtgtgac aaatgtcctc ctggtaccta
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214 ctacacagac agctggcaca ccagtgacga gtgtctatac tgcagccccg tgtgcaagga
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240 cagtttgtgg cgaataaaaa atggcgacca agacaccttg aagggcctaa tgcacgcact
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242 aaagcactca aagacgtacc actttcccaa aactgtcact cagagtctaa agaagaccat
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246 aggtaaccag gtccaatcag taaaaataag ctgcttataa ctggaaatgg ccattgagct
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254 <213> ORGANISM: Homo sapiens
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266 Glu Glu Thr Ser His Gln Leu Leu Cys Asp Lys Cys Pro Pro Gly Thr
270 Tyr Leu Lys Gln His Cys Thr Ala Lys Trp Lys Thr Val Cys Ala Pro
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271	50					55					60				
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278 Leu	Tyr	Cys	Ser		Val	Cys	Lys	Glu		Gln	Tyr	Val	Lys		Glu
279			mı	85	_	_		_	90	_	_	- 1	- 1	95	_
282 Cys 283	Asn	Arg		His	Asn	Arg	vai	Cys 105	GIu	Cys	ьуs	GLu	_	Arg	Tyr
286 Leu	Clu	Tlo	100	Dho	Cvc	LOU	T 1/10		71 **	802	Cvrc	Dro	110	C1,,	Dho
287	GIU	115	Gru	rne	Cys	ьeu	120	птъ	Arg	ser	Cys	125	PIO	сту	Pile
290 Gly	Val		Gln	Δla	Glv	Thr	-	Glu	Ara	Asn	Thr		Cvs	T.vs	Ara
291	130	• • • •	0111	711.0	OLY	135	110	Olu	**** 9	11011	140	•41	Cys	цуо	1119
294 Cys		Asp	Gly	Phe	Phe		Asn	Glu	Thr	Ser		Lys	Ala	Pro	Cys
295 145		-	_		150					155					160
298 Arg	Lys	His	Thr	Asn	Cys	Ser	Val	Phe	Gly	Leu	Leu	Leu	Thr	Gln	Lys
299				165					170					175	
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303			180					185					190		
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307		195	_	_,	_	_,	200	_	_	_	_	205		_	
310 Phe		Val	Pro	Thr	ьys		Thr	Pro	Asn	Trp		Ser	Val	Leu	Val
311	210	Ton	Dro	C1	mh~	215	17-1	7.00	71.7	C1	220	17-1	C1	7\ ~~ ~	Tlo
314 Asp 315 225		ьeu	PIO	Gry	230	ту	Vai	ASII	AIA	235	ser	vai	GIU	Arg	240
318 Lys		Gln	His	Ser		Gln	Glu	Gln	Thr		Gln	T.e.11	T.e.ii	T.vs	
319	9	01		245	001	0111	Olu	0111	250	1110	0111	пси	шец	255	пси
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323	_		260		-			265			-	-	270		
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330 Asn		Thr	Phe	Glu	Gln		Arg	Ser	Leu	Met		Ser	Leu	Pro	Gly
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334 Lys		vaı	GTA	Ата		Asp	тте	GIu	ьуs		тте	ьуs	Ата	Cys	_
335 305 338 Pro		λcr	Cln	Tlo	310	Tuc	T 011	T 011	Sor.	315	Ф~~	7~~	Tlo	Tura	320
339	Det	тэр	GIII	325	пеп	цуз	neu	neu	330	цец	тър	Arg	116	335	ASII
342 Gly	Asp	Gln	Asp		Leu	Lvs	Glv	Leu		His	Ala	Len	Lvs		Ser
343		V	340			J -	011	345		0			350		501
346 Lys	Thr	Tyr	His	Phe	Pro	Lys	Thr		Thr	Gln	Ser	Leu	Lys	Lys	Thr
347		355				-	360					365	_	-	
350 Ile	Arg	Phe	Leu	His	Ser	Phe	Thr	Met	Tyr	Lys	Leu	Tyr	Gln	Lys	Leu
351	370					375					380				
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VERIFICATION SUMMARY

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